	Lys	s Arç	g Glu	900		Gln	ı Lev	ı Glu	905		ı Ile	· Val	. Tyr	910		ı Ala
5	Lys	Glu	915	Val	. Asp	Ala	Leu	920		Asn	Ser	Gln	Tyr 925		Arg	Leu
	Gln	Val 930	Asp	Thr	Asn	Ile	Ala 935		Ile	His	Ala	Ala 940		Lys	Arg	Val
10	His 945	Arg	Ile	Arg	Glu	Ala 950	Tyr	Leu	Pro	Glu	Leu 955	Ser	Val	Ile	Pro	Gly 960
15	Val	Asn	Ala	Ala	Ile 965	Phe	Glu	Glu	Leu	Glu 970	Gly	Arg	Ile	Phe	Thr 975	Ala
13	Tyr	Ser	Leu	Tyr 980	Asp	Ala	Arg	Asn	Val 985	Ile	Lys	Asn	Gly	Asp 990	Phe	Asn
20 ()	Asn	Gly	Leu 995	Leu	Cys	Trp	Asn	Val 1000		Gly	His	Val	Asp		Glu	Glu
The state of the s	Gln	Asn 1010		His	Arg	Ser	Val 1015		Val	Ile	Pro	Glu 1020		Glu	Ala	Glu
25	Val 1025	Ser	Gln	Glu	Val	Arg 1030		Cys	Pro	Gly	Arg 1035		Tyr	Ile	Leu	Arg 1040
3 0	Val	Thr	Ala	Tyr	Lys 1045		Gly	Tyr	Gly	Glu 1050		Cys	Val		Ile 1055	
	Glu	Ile	Glu	Asp 1060	Asn	Thr	Asp	Glu	Leu 1065		Phe	Ser	Asn	Cys 1070	Val	Glu
35	Glu	Glu	Val 1075	Tyr	Pro	Asn	Asn	Thr 1080		Thr	Cys	Asn	Asn 1085		Thr	Gly
	Thr	Gln 1090	Glu	Glu	Tyr		Gly 1095		Tyr	Thr	Ser	Arg 1100		Gln	Gly	Tyr
40	Asp 1105		Ala	Tyr	Gly .	Asn 1110		Pro	Ser		Pro 1115		Asp	Tyr .	Ala	Ser 1120
45	Val	Tyr	Glu		Lys 1125	Ser	Tyr	Thr		Gly 1130		Arg	Glu		Pro 1135	_
	Glu	Ser		Arg 1140	Gly '	Tyr	Gly		Tyr 1145		Pro	Leu		Ala 1150	Gly	Tyr
50	Val		Lys 1155		Leu (Glu		Phe 1160		Glu	Thr		Lys 1165		Trp	Ile
	Glu	Ile	Gly	Glu	Thr	Glu	Gly	Thr	Phe	Ile	Val	Asp	Ser	Val	Glu	Leu

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- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 47 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:62:
- . 15 CGGGGATTAA ATAATTTACC GGCTAGCACG TATCAAGATT GGATAAC
 - (2) INFORMATION FOR SEQ ID NO:63:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 45 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:63:
 - CGGGGATTAA ATAATTTACC GAAAAACGTA TCAAGATTGG ATAAC
 - (2) INFORMATION FOR SEQ ID NO:64:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 23 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:64:
- 40 GGATAGCACT CATCAAAGGT ACC
 - (2) INFORMATION FOR SEQ ID NO:65:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 45 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
- 50 (D) TOPOLOGY: linear
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:65:

CGGGGATTAA ATAATACCGA AAAGCACGTA TCAAGATTGG ATAAC

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	(2) INFORMATION FOR SEQ ID NO:66:	
5	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 45 base pairs(B) TYPE: nucleic acid(C) STRANDEDNESS: single	
10	(D) TOPOLOGY: linear (xi) SEQUENCE DESCRIPTION: SEQ ID NO:66:	
	CGGGGATTAA ATAATTTAAA AAAGCACGTA TCAAGATTGG ATAAC	49
15	(2) INFORMATION FOR SEQ ID NO:67:	
20	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 45 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
25	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:67:	
Moore Bear	CGGGGATTAA ATAATTTACC GAAGCACGTA TCAAGATTGG ATAAC	45
30	(2) INFORMATION FOR SEQ ID NO:68:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 51 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:68:	
40	GGATTAAATA ATTTACCGAA AAGCATATCA AGATTGGATA ACATATAATC G	51
	(2) INFORMATION FOR SEQ ID NO:69:	
45	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 51 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
50	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:69:	
	GGATTAAATA ATTTACCGAA AAGCACGACA AGATTGGATA ACATATAATC C	E 1